SEQUENCE LISTING

<110>   Liaw, Gin
        Pedersen, Sven
        Hendriksen, Sven

<120>   A Method of Producing Saccharide Preparations

<130>   5318.200-US

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<170>   PatentIn version 3.2

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ttg gca aat gtg att tcc aag cgc gcg acc ttg gat tca tgg ttg agc   96
Leu Ala Asn Val Ile Ser Lys Arg Ala Thr Leu Asp Ser Trp Leu Ser
            -5   -1 1   5

aac gaa gcg acc gtg gct cgt act gcc atc ctg aat aac atc ggg gcg   144
Asn Glu Ala Thr Val Ala Arg Thr Ala Ile Leu Asn Asn Ile Gly Ala
    10   15   20

gac ggt gct tgg gtg tcg ggc gcg gac tct ggc att gtc gtt gct agt   192
Asp Gly Ala Trp Val Ser Gly Ala Asp Ser Gly Ile Val Val Ala Ser
25   30   35   40

ccc agc acg gat aac ccg gac tac ttc tac acc tgg act cgc gac tct   240
Pro Ser Thr Asp Asn Pro Asp Tyr Phe Tyr Thr Trp Thr Arg Asp Ser
                45   50   55

ggt ctc gtc ctc aag acc ctc gtc gat ctc ttc cga aat gga gat acc   288
Gly Leu Val Leu Lys Thr Leu Val Asp Leu Phe Arg Asn Gly Asp Thr
            60   65   70

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agt ctc ctc tcc acc att gag aac tac atc tcc gcc cag gca att gtc    336
Ser Leu Leu Ser Thr Ile Glu Asn Tyr Ile Ser Ala Gln Ala Ile Val
        75                  80                  85

cag ggt atc agt aac ccc tct ggt gat ctg tcc agc ggc gct ggt ctc    384
Gln Gly Ile Ser Asn Pro Ser Gly Asp Leu Ser Ser Gly Ala Gly Leu
        90                  95                  100

ggt gaa ccc aag ttc aat gtc gat gag act gcc tac act ggt tct tgg    432
Gly Glu Pro Lys Phe Asn Val Asp Glu Thr Ala Tyr Thr Gly Ser Trp
105                 110                 115                 120

gga cgg ccg cag cga gat ggt ccg gct ctg aga gca act gct atg atc    480
Gly Arg Pro Gln Arg Asp Gly Pro Ala Leu Arg Ala Thr Ala Met Ile
                125                 130                 135

ggc ttc ggg cag tgg ctg ctt gac aat ggc tac acc agc acc gca acg    528
Gly Phe Gly Gln Trp Leu Leu Asp Asn Gly Tyr Thr Ser Thr Ala Thr
                140                 145                 150

gac att gtt tgg ccc ctc gtt agg aac gac ctg tcg tat gtg gct caa    576
Asp Ile Val Trp Pro Leu Val Arg Asn Asp Leu Ser Tyr Val Ala Gln
        155                 160                 165

tac tgg aac cag aca gga tat gat ctc tgg gaa gaa gtc aat ggc tcg    624
Tyr Trp Asn Gln Thr Gly Tyr Asp Leu Trp Glu Glu Val Asn Gly Ser
        170                 175                 180

tct ttc ttt acg att gct gtg caa cac cgc gcc ctt gtc gaa ggt agt    672
Ser Phe Phe Thr Ile Ala Val Gln His Arg Ala Leu Val Glu Gly Ser
185                 190                 195                 200

gcc ttc gcg acg gcc gtc ggc tcg tcc tgc tcc tgg tgt gat tct cag    720
Ala Phe Ala Thr Ala Val Gly Ser Ser Cys Ser Trp Cys Asp Ser Gln
                205                 210                 215

gca ccc gaa att ctc tgc tac ctg cag tcc ttc tgg acc ggc agc ttc    768
Ala Pro Glu Ile Leu Cys Tyr Leu Gln Ser Phe Trp Thr Gly Ser Phe
                220                 225                 230

att ctg gcc aac ttc gat agc agc cgt tcc ggc aag gac gca aac acc    816
Ile Leu Ala Asn Phe Asp Ser Ser Arg Ser Gly Lys Asp Ala Asn Thr
                235                 240                 245

ctc ctg gga agc atc cac acc ttt gat cct gag gcc gca tgc gac gac    864
Leu Leu Gly Ser Ile His Thr Phe Asp Pro Glu Ala Ala Cys Asp Asp
        250                 255                 260

tcc acc ttc cag ccc tgc tcc ccg cgc gcg ctc gcc aac cac aag gag    912
Ser Thr Phe Gln Pro Cys Ser Pro Arg Ala Leu Ala Asn His Lys Glu
265                 270                 275                 280

gtt gta gac tct ttc cgc tca atc tat acc ctc aac gat ggt ctc agt    960
Val Val Asp Ser Phe Arg Ser Ile Tyr Thr Leu Asn Asp Gly Leu Ser
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gac agc gag gct gtt gcg gtg ggt cgg tac cct gag gac acg tac tac    1008
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2

Asp Ser Glu Ala Val Ala Val Gly Arg Tyr Pro Glu Asp Thr Tyr Tyr
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aac ggc aac ccg tgg ttc ctg tgc acc ttg gct gcc gca gag cag ttg   1056
Asn Gly Asn Pro Trp Phe Leu Cys Thr Leu Ala Ala Ala Glu Gln Leu
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tac gat gct cta tac cag tgg gac aag cag ggg tcg ttg gag gtc aca   1104
Tyr Asp Ala Leu Tyr Gln Trp Asp Lys Gln Gly Ser Leu Glu Val Thr
        330   335   340

gat gtg tcg ctg gac ttc ttc aag gca ctg tac agc gat gct gct act   1152
Asp Val Ser Leu Asp Phe Phe Lys Ala Leu Tyr Ser Asp Ala Ala Thr
345   350   355   360

ggc acc tac tct tcg tcc agt tcg act tat agt agc att gta gat gcc   1200
Gly Thr Tyr Ser Ser Ser Ser Ser Thr Tyr Ser Ser Ile Val Asp Ala
                365   370   375

gtg aag act ttc gcc gat ggc ttc gtc tct att gtg gaa act cac gcc   1248
Val Lys Thr Phe Ala Asp Gly Phe Val Ser Ile Val Glu Thr His Ala
                380   385   390

gca agc aac ggc tcc atg tcc gag caa tac gac aag tct gat ggc gag   1296
Ala Ser Asn Gly Ser Met Ser Glu Gln Tyr Asp Lys Ser Asp Gly Glu
        395   400   405

cag ctt tcc gct cgc gac ctg acc tgg tct tat gct gct ctg ctg acc   1344
Gln Leu Ser Ala Arg Asp Leu Thr Trp Ser Tyr Ala Ala Leu Leu Thr
        410   415   420

gcc aac aac cgt cgt aac tcc gtc gtg cct gct tct tgg ggc gag acc   1392
Ala Asn Asn Arg Arg Asn Ser Val Val Pro Ala Ser Trp Gly Glu Thr
425   430   435   440

tct gcc agc agc gtg ccc ggc acc tgt gcg gcc aca tct gcc att ggt   1440
Ser Ala Ser Ser Val Pro Gly Thr Cys Ala Ala Thr Ser Ala Ile Gly
                445   450   455

acc tac agc agt gtg act gtc acc tcg tgg ccg agt atc gtg gct act   1488
Thr Tyr Ser Ser Val Thr Val Thr Ser Trp Pro Ser Ile Val Ala Thr
                460   465   470

ggc ggc acc act acg acg gct acc ccc act gga tcc ggc agc gtg acc   1536
Gly Gly Thr Thr Thr Thr Ala Thr Pro Thr Gly Ser Gly Ser Val Thr
                475   480   485

tcg acc agc aag acc acc gcg act gct agc aag acc agc acc acg acc   1584
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cgc tct ggt atg tca ctg tga   1605
Arg Ser Gly Met Ser Leu
505   510

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Asp Gly Ala Trp Val Ser Gly Ala Asp Ser Gly Ile Val Val Ala Ser
25   30   35   40

Pro Ser Thr Asp Asn Pro Asp Tyr Phe Tyr Thr Trp Thr Arg Asp Ser
             45   50   55

Gly Leu Val Leu Lys Thr Leu Val Asp Leu Phe Arg Asn Gly Asp Thr
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Ser Leu Leu Ser Thr Ile Glu Asn Tyr Ile Ser Ala Gln Ala Ile Val
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Gln Gly Ile Ser Asn Pro Ser Gly Asp Leu Ser Ser Gly Ala Gly Leu
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Gly Glu Pro Lys Phe Asn Val Asp Glu Thr Ala Tyr Thr Gly Ser Trp
105   110   115   120

Gly Arg Pro Gln Arg Asp Gly Pro Ala Leu Arg Ala Thr Ala Met Ile
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Gly Phe Gly Gln Trp Leu Leu Asp Asn Gly Tyr Thr Ser Thr Ala Thr
             140   145   150

Asp Ile Val Trp Pro Leu Val Arg Asn Asp Leu Ser Tyr Val Ala Gln
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Tyr Trp Asn Gln Thr Gly Tyr Asp Leu Trp Glu Glu Val Asn Gly Ser
     170   175   180

4

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Ser Phe Phe Thr Ile Ala Val Gln His Arg Ala Leu Val Glu Gly Ser
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Ala Phe Ala Thr Ala Val Gly Ser Ser Cys Ser Trp Cys Asp Ser Gln
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Ala Pro Glu Ile Leu Cys Tyr Leu Gln Ser Phe Trp Thr Gly Ser Phe
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Ile Leu Ala Asn Phe Asp Ser Ser Arg Ser Gly Lys Asp Ala Asn Thr
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Leu Leu Gly Ser Ile His Thr Phe Asp Pro Glu Ala Ala Cys Asp Asp
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Ser Thr Phe Gln Pro Cys Ser Pro Arg Ala Leu Ala Asn His Lys Glu
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Val Val Asp Ser Phe Arg Ser Ile Tyr Thr Leu Asn Asp Gly Leu Ser
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Asp Ser Glu Ala Val Ala Val Gly Arg Tyr Pro Glu Asp Thr Tyr Tyr
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Asn Gly Asn Pro Trp Phe Leu Cys Thr Leu Ala Ala Ala Glu Gln Leu
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Tyr Asp Ala Leu Tyr Gln Trp Asp Lys Gln Gly Ser Leu Glu Val Thr
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Asp Val Ser Leu Asp Phe Phe Lys Ala Leu Tyr Ser Asp Ala Ala Thr
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Gly Thr Tyr Ser Ser Ser Ser Ser Thr Tyr Ser Ser Ile Val Asp Ala
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Val Lys Thr Phe Ala Asp Gly Phe Val Ser Ile Val Glu Thr His Ala
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Ala Ser Asn Gly Ser Met Ser Glu Gln Tyr Asp Lys Ser Asp Gly Glu
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Gln Leu Ser Ala Arg Asp Leu Thr Trp Ser Tyr Ala Ala Leu Leu Thr
    410             415             420


Ala Asn Asn Arg Arg Asn Ser Val Val Pro Ala Ser Trp Gly Glu Thr
425             430             435             440


Ser Ala Ser Ser Val Pro Gly Thr Cys Ala Ala Thr Ser Ala Ile Gly
            445             450             455


Thr Tyr Ser Ser Val Thr Val Thr Ser Trp Pro Ser Ile Val Ala Thr
        460             465             470


Gly Gly Thr Thr Thr Thr Ala Thr Pro Thr Gly Ser Gly Ser Val Thr
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Ser Thr Ser Lys Thr Thr Ala Thr Ala Ser Lys Thr Ser Thr Thr Thr
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Arg Ser Gly Met Ser Leu
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